

L Number	Hits	Search Text	DB	Time stamp
3	0	ephrin adj A adj "8"	USPAT; US-PGPUB; EPO; JPO; DERWENT	2003/12/11 14:32
4	36	ephrin SAME receptor SAME "8"	USPAT; US-PGPUB; EPO; JPO; DERWENT	2003/12/11 14:33
5	24	human SAME eek	USPAT; US-PGPUB; EPO; JPO; DERWENT	2003/12/11 14:33
6	59	(ephrin SAME receptor SAME "8") OR (human SAME eek)	USPAT; US-PGPUB; EPO; JPO; DERWENT	2003/12/11 14:49
7	218	Prayaga.IN. OR Taupier.IN. OR Bandaru.IN.	USPAT; US-PGPUB; EPO; JPO; DERWENT	2003/12/11 14:49
8	4	(Prayaga.IN. OR Taupier.IN. OR Bandaru.IN.) AND ephrin	USPAT; US-PGPUB; EPO; JPO; DERWENT	2003/12/11 14:50

FILE 'MEDLINE, EMBASE, USPATFULL, BIOSIS, CAPLUS' ENTERED AT 14:52:39 ON  
11 DEC 2003

L1	13 S EPHRINA8 OR (EPHRIN A8)
L2	37 S HUMAN (S) EEK
L3	0 S (PRAYAGA OR TAUPIER OR BANDARU) AND EPHRIN
L4	4 S (PARAYAGA OR TAUPIER OR BANDARU) AND NOV
L5	54 S L1 OR L2 OR L4
L6	48 DUP REM L5 (6 DUPLICATES REMOVED)

Query Match 100.0%; Score 2744; DB 1; Length 1005;  
Best Local Similarity 100.0%; Pred. No. 3e-213;  
Matches 515; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy	1	ARGEVNLLDTSTIHGDWGLTYPAHGWDSINEVDES	FQPIHTYQVCNVMSPNQNNWLRTS	60
Db	28	ARGEVNLLDTSTIHGDWGLTYPAHGWDSINEVDES	FQPIHTYQVCNVMSPNQNNWLRTS	87
Qy	61	WVPRDGARRVYAEIKFTLRDCNSMPGVLGTCKET	FNLYYLES	120
Db	88	WVPRDGARRVYAEIKFTLRDCNSMPGVLGTCKET	FNLYYLES	147
Qy	121	IAADESFTGADLGVRRLKLNTEVRSVGPLSKRG	FYLA	180
Db	148	IAADESFTGADLGVRRLKLNTEVRSVGPLSKRG	FYLA	207
Qy	181	VRNLAAFSEAVTGADSSSLVEVRGQCVRHSEER	DTPKMYCSAEGEWLVPIGKCVCSAGYE	240
Db	208	VRNLAAFSEAVTGADSSSLVEVRGQCVRHSEER	DTPKMYCSAEGEWLVPIGKCVCSAGYE	267
Qy	241	ERRDACVACELGFYKSAPGDQLCARCPPHSHSA	APAAQACHCDLSYYRAALDPPSSACTR	300
Db	268	ERRDACVACELGFYKSAPGDQLCARCPPHSHSA	APAAQACHCDLSYYRAALDPPSSACTR	327
Qy	301	PPSAPVNLISSVNGTSVTLEWAPPLDPGGRSDI	TYN	360
Db	328	PPSAPVNLISSVNGTSVTLEWAPPLDPGGRSDI	TYN	387
Qy	361	QQTSLVQASLLVANLLAHMNYSFWIEAVNGVSD	LSPEPRRAAVVNITTNQAAPSQVVVIR	420
Db	388	QQTSLVQASLLVANLLAHMNYSFWIEAVNGVSD	LSPEPRRAAVVNITTNQAAPSQVVVIR	447
Qy	421	QERAGQTSVSLWQEPEQPNGIILEYEIKYYEKD	KEMQSYSTLKAVTTRATVSGLKPGTR	480
Db	448	QERAGQTSVSLWQEPEQPNGIILEYEIKYYEKD	KEMQSYSTLKAVTTRATVSGLKPGTR	507
Qy	481	YVFQVRARTSAGCGRFSQAMEVETGKPRPRYD	TRT	515
Db	508	YVFQVRARTSAGCGRFSQAMEVETGKPRPRYD	TRT	542